HT001

wt ORF (SEQ ID NO: 1)

MQRPNAHRISQPIRQIÍYGLLNASPHLDKTSWNALPPQPLAFSEVERINKNIRTSIIDAVELAKDHSDLSRLTELSLRRRQMLLLETLKV KQTILEPIPTSLKLPIAVSCYWLQHTETKAKLHHLQSLLTMLVGPLIAIINSPGKEELQEDGAKMLYAEFQRVKAQTRLGTRLDLDTAHI FCQWQSCLQMGMYLNQLLSTPLPEPDLTRLYSGSLVHGLCQQLLASTSVESLLSICPEAKQLYEYLFNATRSYAPAEIFLPKGRSNSK KKRQKKQNTSCSKNRGRTTAHTKCWYEGNNRFGLLMVENLEEHSEASNIE

(-1) ORF (SEQ ID NO: 2)

MÓRPNAHRISQPIRQIIÝGLLNASPHLDKTSWNALPPQPLAFSEVERINKNIRTSIIDAVELAKDHSDLSRLTELSLRRRQMLLLETLKV KQTILEPIPTSLKLPIAVSCYWLQHTETKAKLHHLQSLLLTMLVGPLIAIINSPGKEELQEDGAKMLYAEFQRVKAQTRLGTRLDLDTAHI FCQWQSCLQMGMYLNQLLSTPLPEPDLTRLYSGSLVHGLCQQLLASTSVESLLSICPEAKQLYEYLFNATRSYAPAEIFLPKGRSNSK KKGRRNRIPAVLRTEGEPLHTPSVGMRETTGLGC

(+1)/(-2) ORF (SEQ ID NO: 3/118)

MQRPNAHRISQPIRQIIYGLLNASPHLDKTSWNALPPQPLAFSEVERINKNIRTSIIDAVELAKDHSDLSRLTELSLRRRQMLLLETLKV KQTILEPIPTSLKLPIAVSCYWLQHTETKAKLHHLQSLLLTMLVGPLIAIINSPGKEELQEDGAKMLYAEFQRVKAQTRLGTRLDLDTAHI FCQWQSCLQMGMYLNQLLSTPLPEPDLTRLYSGSLVHGLCQQLLASTSVESLLSICPEAKQLYEYLFNATRSYAPAEIFLPKGRSNSK KK/K/AEETEYQLE

U79260

wt ORF (SEQ ID NO: 4)

MGHPRÁIQPSVFFSPÝDVHFLLYPIRCPYLKIGRFHIKLKGLHFLFSFLFFFETQSHSVTRLECSGTISAHCNLCLPGSSNSPASASRV AGTAGTCRRAQLIFVFLAEMGFHHVGRDGLDLNLVIHPPRSPKALGLQA

(-1)ORF (SEQ ID NO: 5)

MGHPRÄIQPSVFFSPYDVHFLLYPIRCPYLKIGRFHIKLKGLHFLFSFLFFF<u>LRHSLTLSPGWSAVARSRLTATSASQVQVILLPQPPEW</u> <u>LGLQARAAAPS</u>

(+1)/(-2)ORF (SEQ ID NO: 6)

MGHPRAIQPSVFFSPYDVHFLLYPIRCPYLKIGRFHIKLKGLHFLFSFLFFF(F)

PTHL3

(wt)ORF (SEQ ID NO: 7)

MQRRLVQQWSVAVFLLSYAVPSCGRSVEGLSRRLKRAVSEHQLLHDKGKSIQDLRRRFFLHHLIAEIHTAEIRATSEVSPNSKPSPNT KNHPVRFGSDDEGRYLTQETNKVETYKEQPLKTPGKKKKGKPGKRKEQEKKKRRTRSAWLDSGVTGSGLEGDHLSDTSTTSLELD SRTALLWGLKKKKENNRRTHHMQLMISLFKSPLLLL

(-1)ORF (SEQ ID NO: 8)

MÓRRLVQQWSVAVFLLSYAVPSCGRSVEGLSRRLKRAVSEHQLLHDKGKSIQDLRRRFFLHHLIAEIHTAEIRATSEVSPNSKPSPNT KNHPVRFGSDDEGRYLTQETNKVETYKEQPLKTPGKKKKGKPGKRKEQEKKKRRTRSAWLDSGVTGSGLEGDHLSDTSTTSLELD SRTALLWGLKKK*RKTTEEHIICN*

(+1)/(-2)ORF (SEQ ID NO: 9)

MQRRLVQQWSVAVFLLSYAVPSCGRSVEGLSRRLKRAVSEHQLLHDKGKSIQDLRRRFFLHHLIAEIHTAEIRATSEVSPNSKPSPNT KNHPVRFGSDDEGRYLTQETNKVETYKEQPLKTPGKKKKGKPGKRKEQEKKKRRTRSAWLDSGVTGSGLEGDHLSDTSTTSLELD SRTALLWGLKKK(K)GKQQKNTSYATNDLII

TGFbRII

(wt) (SEQ ID NO: 10)

MGRGLLRGLWPLHÍVLWTRIASTIPPHVQKSVNNDMIVTDNNGAVKFPQLCKFCDVRFSTCDNQKSCMSNCSITSICEKPQEVCVAV WRKNDENITLETVCHDPKLPYHDFILEDAASPKCIMKEKKKPGETFFMCSCSSDECNDNIIFSEEYNTSNPDLLLVIFQVTGISLLPPLG VAISVIIIFYCYRVNRQQKLSSTWETGKTRKLMEFSEHCAIILEDDRSDISSTCANNINHNTELLPIELDTLVGKGRFAEVYKAKLKQNTS EQFETVAVKIFPYEEYASWKTEKDIFSDINLKHENILQFLTAEERKTELGKQYWLITAFHAKGNLQEYLTRHVISWEDLRKLGSSLARGI AHLHSDHTPCGRPKMPIVHRDLNSSNILVKNDLTCCLCDFGLSLRLDPTLSVDDLANSGQVGTARYMAPEVLESRMNLENAESFKQT DVYSMALVLWEMTSRCNAVGEVKDYEPPFGSKVREHPCVESMKDNVLRDRGRPEIPSFWLNHQGIQMVCETLTECWDHDPEARLT AQCVAERFSELEHLDRLSGRSCSEEKIPEDGSLNTTK

(-1)ORF (SEQ ID NO: 11)

MGRGLLRGLWPLHIVLWTRIASTIPPHVQKSVNNDMIVTDNNGAVKFPQLCKFCDVRFSTCDNQKSCMSNCSITSICEKPQEVCVAV WRKNDENITLETVCHDPKLPYHDFILEDAASPKCIMKEKK<u>SLVRLSSCVPVALMSAMTTSSSQKNITPAILTCC</u>

(+1)/(-2)ORF (SEQ ID NO: 12/119)

MGRGLLRGLWPLHIVLWTRIASTIPPHVQKSVNNDMIVTDNNGAVKFPQLCKFCDVRFSTCDNQKSCMSNCSITSICEKPQEVCVAV WRKNDENITLETVCHDPKLPYHDFILEDAASPKCIMKEKK(K) \underline{AW}

MACS

(wt)ORF (SEQ ID NO: 13)

MGAQFSKTAAKGEAAAÉRPGEAAVASSPSKANGQENGHVKVNGDASPAAAESGAKEELQANGSAPAADKEEPAAAGSGAASPSS AEKGEPAAAAAPEAGASPVEKEAPAEGEAAEPGSATAAEGEAASAASSTSSPKAEDGATPSPSNETPKKKKKRFSFKKSFKLSGFS FKKNKKEAGEGGEAEAPAAEGGKDEAAGGAAAAAAEAGAASGEQAAAPGEEAAAGEEGAAGGDPQEAKPQEAAVAPEKPPASDE TKAAEEPSKVEEKKAEEAGASAAACEAPSAAGPGAPPEQEAAPAEEPAAAAASSACAAPSQEAQPECSPEAPPAEAAE

(-1)ORF (SEQ ID NO: 14)

MGAQFSKTAAKGEAAAERPGEAAVASSPSKANGQENGHVKVNGDASPAAAESGAKEELQANGSAPAADKEEPAAAGSGAASPSS AEKGEPAAAAAPEAGASPVEKEAPAEGEAAEPGSATAAEGEAASAASSTSSPKAEDGATPSPSNETPKKK<u>RSAFPSRSLSS</u>

(+1)/(-2)ORF (SEQ ID NO: 15)

MGAQFSKTAAKGEAAAERPGEAAVASSPSKANGQENGHVKVNGDASPAAAESGAKEELQANGSAPAADKEEPAAAGSGAASPSS AEKGEPAAAAAPEAGASPVEKEAPAEGEAAEPGSATAAEGEAASAASSTSSPKAEDGATPSPSNETPKKK(K)<u>EALFLQEVFQAERL</u> **LLQEEQEGGWRRR**

TCF-4

(wt)ORF (SEQ ID NO: 16)

MPQLNGGGGDDLGANDELISFKDEGEQEEKSSENSSAERDLADVKSSLVNESETNQNSSSDSEAERRPPPRSESFRDKSRESLEEA AKRQDGGLFKGPPYPGYPFIMIPDLTSPYLPNGSLSPTARTYLQMKWPLLDVQAGSLQSRQALKDARSPSPAHIVSNKVPVVQHPHH VHPLTPLITYSNEHFTPGNPPPHLPADVDPKTGIPRPPHPPDISPYYPLSPGTVGQIPHPLGWLVPQQGQPVYPITTGGFRHPYPTALT VNASVSRFPPHMVPPHHTLHTTGIPHPAIVTPTVKQESSQSDVGSLHSSKHQDSKKEEEKKKPHIKKPLNAFMLYMKEMRAKVVAEC TLKESAAINQILGRRWHALSREEQAKYYELARKERQLHMQLYPGWSARDNYGKKKKRKDKQPGETNEHSECFLNPCLSLPPITDLS APKKCRARFGLDQQNNWCGPCRRKKKCVRYIQGEGSCLSPPSSDGSLLDSPPPSPNLLGSPPRDAKSQTEQTQPLSLSLKPDPLAH LSMMPPPPALLLAEATHKASALCPNGALDLPPAALQPAAPSSSIAQPSTSWLHSHSSLAGTQPQPLSLVTKSLE

(-1)ORF (SEQ ID NO: 17)

MPQLNGGGGDDLGANDELISFKDEGEQEEKSSENSSAERDLADVKSSLVNESETNQNSSSDSEAERRPPPRSESFRDKSRESLEEA AKRQDGGLFKGPPYPGYPFIMIPDLTSPYLPNGSLSPTARTYLQMKWPLLDVQAGSLQSRQALKDARSPSPAHIVSNKVPVVQHPHH VHPLTPLITYSNEHFTPGNPPPHLPADVDPKTGIPRPPHPPDISPYYPLSPGTVGQIPHPLGWLVPQQGQPVYPITTGGFRHPYPTALT VNASVSRFPPHMVPPHHTLHTTGIPHPAIVTPTVKQESSQSDVGSLHSSKHQDSKKEEEKKKPHIKKPLNAFMLYMKEMRAKVVAEC TLKESAAINQILGRRWHALSREEQAKYYELARKERQLHMQLYPGWSARDNYGKKKKRKRDKQPGETNEHSECFLNPCLSLPPITDLS APKKCRARFGLDQQNNWCGPCRRKKSAFATYKVKAAASAHPLQMEAY

(+1)/(-2)ORF (SEQ ID NO: 18)

MPQLNGGGGDDLGANDELISFKDEGEQEEKSSENSSAERDLADVKSSLVNESETNQNSSSDSEAERRPPPRSESFRDKSRESLEEA AKRQDGGLFKGPPYPGYPFIMIPDLTSPYLPNGSLSPTARTYLQMKWPLLDVQAGSLQSRQALKDARSPSPAHIVSNKVPVVQHPHH VHPLTPLITYSNEHFTPGNPPPHLPADVDPKTGIPRPPHPPDISPYYPLSPGTVGQIPHPLGWLVPQQGQPVYPITTGGFRHPYPTALT VNASVSRFPPHMVPPHHTLHTTGIPHPAIVTPTVKQESSQSDVGSLHSSKHQDSKKEEEKKKPHIKKPLNAFMLYMKEMRAKVVAEC TLKESAAINQILGRRWHALSREEQAKYYELARKERQLHMQLYPGWSARDNYGKKKKRKDKQPGETNEHSECFLNPCLSLPPITDLS APKKCRARFGLDQQNNWCGPCRRKK(K)VRSLHTR

(wt)ORF (SEQ ID NO: 19)

IPAFPAGTVLQPFPEAALATRVTVPAVEAPAAPRLDLEESEEFKERCTQCAAVSWGLTDEGKYYCTSCHNVTERYQEVTNTDLIPNT QIKALNRGLKKKNNTEKGWDWYVCEGFQYILYQQAEALKNLGVGPELKNDVLHNFWKRYLQKSKQAYCKNPVYTTGRKPTVLEDNL SHSDWASEPELLSDVSCPPFLESGAESQSDIHTRKPFPVSKASQSETSVCSGSLDGVEYSQRKEKGIVKMTMPQTLAFCYLSLLWQ REAITLSDLLRFVEEDHIPYINAFQHFPEQMKLYGRDRGIFGIESWPDYEDIYKKTIEVGTFLDLPRFPDITEDCYLHPNILCMKYLMEVN LPDEMHSLTCHVVKMTGMGEVDFLTFDPIAKMAKAVKYDVQAVAIIVVVLKLLFLMDDSFEWSLSNLAEKHNEKNKKDKPWFDFRKW YQIMKKAFDEKKQKWEEARAKYLWKSEKPLYYSFVDKPVAYKKREMVVNLQKQFSTLVDSTATAGKKSPSSFQFNWTEEDTDRTC FHGHSLQGVLKEKGQSLLTKNSLYWLSTQKFCRW

(-1)ORF (SEQ ID NO: 20)

IPAFPAGTVLQPFPEAALATRVTVPAVEAPAAPRLDLEESEEFKERCTQCAAVSWGLTDEGKYYCTSCHNVTERYQEVTNTDLIPNT QIKALNRGLKKK<u>TILKKAGIGMCVKVSSIFFINKQKP</u>

(+1)/(-2)ORF (SEQ ID NO: 21/120)

IPAFPAGTVLQPFPEAALATRVTVPAVEAPAAPRLDLEESEEFKERCTQCAAVSWGLTDEGKYYCTSCHNVTERYQEVTNTDLIPNT QIKALNRGLKKK(K)QY

AC-1

(wt)ORF (SEQ ID NO: 22)

MDTQKQIHKTHNSKNQFFTIFFFLSVEFGKEGTRKNFYLLLSIGHYGRKSRRADLGTADTADKTEPECFAASWTFDPNPSVTVSGAHS

(-1)ORF (SEQ ID NO: 23)

MDTQKQIHKTHNSKNQFFTIFFSCQLNLGRKEHAKIFTFFQLDTMDGNPGELTLELQTLQIKQSQNALLPAGPLTQTPV

(+1)/(-2)ORF (SEQ ID NO: 24)

MDTQKQIHKTHNSKNQFFTIFF(F)PVS

(wt)ORF (SEQ ID NO: 25)

MAGQQFQYDDSGNTFFYFLTSFVGLIVIPATYYLWPRDQNAEQIRLKNIRKVYGRCMWYRLRLLKPQPNIIPTVKKIVLLAGWALFLFL AYKVSKTDREYQEYNPYEVLNLDPGATVAEIKKQYRLLSLKYHPDKGGDEVMFMRIAKAYAALTDEESRKNWEEFGNPDGPQATSF GIALPAWIVDQKNSILVLLVYGLAFMVILPVVVGSWWYRSIRYSGDQILIRTTQIYTYFVYKTRNMDMKRLIMVLAGASEFDPQYNKDAT SRPTDNILIPQLIREIGSINLKKNEPPLTCPYSLKARVLLLSHLARMKIPETLEEDQQFMLKKCPALLQEMVNVICQLIVMARNREEREFR APTLASLENCMKLSQMAVQGLQQFKSPLLQLPHIEEDNLRRVSNHKKYKIKTIQDLVSLKESDRHTLLHFLEDEKYEEVMAVLGSFPY VTMDIKSQVLDDEDSNNITVGSLVTVLVKLTRQTMAEVFEKEQSICAAEEQPAEDGQGETNKNRTKGGWQQKSKGPKKTAKSKKKK PLKKKPTPVLLPQSKQQKQKQANGVVGNEAAVKEDEEEVSDKGSDSEEEETNRDSQSEKDDGSDRDSDREQDEKQNKDDEAEW QELQQSIQRKERALLETKSKITHPVYSLYFPEEKQEWWWLYIADRKEQTLISMPYHVCTLKDTEEVELKFPAPGKPGNYQYTVFLRSD SYMGLDQIKPLKLEVHEAKPVPENHPQWDTAIEGDEDQEDSEGFEDSFEEEEEEEDDD

(-1) 9er A-Repeat (SEQ ID NO: 26)

MAGQQFQYDDSGNTFFYFLTSFVGLIVIPATYYLWPRDQNAEQIRLKNIRKVYGRCMWYRLRLLKPQPNIIPTVKKIVLLAGWALFLFL AYKVSKTDREYQEYNPYEVLNLDPGATVAEIKKQYRLLSLKYHPDKGGDEVMFMRIAKAYAALTDEESRKNWEEFGNPDGPQATSF GIALPAWIVDQKNSILVLLVYGLAFMVILPVVVGSWWYRSIRYSGDQILIRTTQIYTYFVYKTRNMDMKRLIMVLAGASEFDPQYNKDAT SRPTDNILIPQLIREIGSINLKKNEPPLTCPYSLKARVLLLSHLARMKIPETLEEDQQFMLKKCPALLQEMVNVICQLIVMARNREEREFR APTLASLENCMKLSQMAVQGLQQFKSPLLQLPHIEEDNLRRVSNHKKYKIKTIQDLVSLKESDRHTLLHFLEDEKYEEVMAVLGSFPY VTMDIKSQVLDDEDSNNITVGSLVTVLVKLTRQTMAEVFEKEQSICAAEEQPAEDGQGETNKNRTKGGWQQKSKGPKKTAKSKK<u>RN</u>

(+1)/(-2) 9er A-Repeat (SEQ ID NO: 27)

MAGQQFQYDDSGNTFFYFLTSFVGLIVIPATYYLWPRDQNAEQIRLKNIRKVYGRCMWYRLRLKPQPNIIPTVKKIVLLAGWALFLFL AYKVSKTDREYQEYNPYEVLNLDPGATVAEIKKQYRLLSLKYHPDKGGDEVMFMRIAKAYAALTDEESRKNWEEFGNPDGPQATSF GIALPAWIVDQKNSILVLLVYGLAFMVILPVVVGSWWYRSIRYSGDQILIRTTQIYTYFVYKTRNMDMKRLIMVLAGASEFDPQYNKDAT SRPTDNILIPQLIREIGSINLKKNEPPLTCPYSLKARVLLLSHLARMKIPETLEEDQQFMLKKCPALLQEMVNVICQLIVMARNREEREFR APTLASLENCMKLSQMAVQGLQQFKSPLLQLPHIEEDNLRRVSNHKKYKIKTIQDLVSLKESDRHTLLHFLEDEKYEEVMAVLGSFPY VTMDIKSQVLDDEDSNNITVGSLVTVLVKLTRQTMAEVFEKEQSICAAEEQPAEDGQGETNKNRTKGGWQQ KSKGPKKTAKSKK(K)<u>ETFKKKTYTCAITTVKATETKAGKWSRWE</u>

(-1) 10er A-Repeat (SEQ ID NO: 28)

MÁGQQFQYDDSGNTFFYFLTSFVÖLIVIPATYYLWPRDQNAEQIRLKNIRKVYGRCMWYRLRLKPQPNIIPTVKKIVLLAGWALFLFL AYKVSKTDREYQEYNPYEVLNLDPGATVAEIKKQYRLLSLKYHPDKGGDEVMFMRIAKAYAALTDEESRKNWEEFGNPDGPQATSF GIALPAWIVDQKNSILVLLVYGLAFMVILPVVVGSWWYRSIRYSGDQILIRTTQIYTYFVYKTRNMDMKRLIMVLAGASEFDPQYNKDAT SRPTDNILIPQLIREIGSINLKKNEPPLTCPYSLKARVLLLSHLARMKIPETLEEDQQFMLKKCPALLQEMVNVICQLIVMARNREEREFR APTLASLENCMKLSQMAVQGLQQFKSPLLQLPHIEEDNLRRVSNHKKYKIKTIQDLVSLKESDRHTLLHFLEDEKYEEVMAVLGSFPY VTMDIKSQVLDDEDSNNITVGSLVTVLVKLTRQTMAEVFEKEQSICAAEEQPAEDGQGETNKNRTKGGWQQ KSKGPKKTAKSKKKKPLKK*NLHLCYYHSQSNRNKSRQMESLGMKLQ*

(+1)/(-2) 10er A-Repeat (SEQ ID NO: 29)

MAGQQFQYDDSGNTFFYFLTSFVGLIVIPATYYLWPRDQNAEQIRLKNIRKVYGRCMWYRLRLLKPQPNIIPTVKKIVLLAGWALFLFL AYKVSKTDREYQEYNPYEVLNLDPGATVAEIKKQYRLLSLKYHPDKGGDEVMFMRIAKAYAALTDEESRKNWEEFGNPDGPQATSF GIALPAWIVDQKNSILVLLVYGLAFMVILPVVVGSWWYRSIRYSGDQILIRTTQIYTYFVYKTRNMDMKRLIMVLAGASEFDPQYNKDAT SRPTDNILIPQLIREIGSINLKKNEPPLTCPYSLKARVLLLSHLARMKIPETLEEDQQFMLKKCPALLQEMVNVICQLIVMARNREEREFR APTLASLENCMKLSQMAVQGLQQFKSPLLQLPHIEEDNLRRVSNHKKYKIKTIQDLVSLKESDRHTLLHFLEDEKYEEVMAVLGSFPY VTMDIKSQVLDDEDSNNITVGSLVTVLVKLTRQTMAEVFEKEQSICAAEEQPAEDGQGETNKNRTKGGWQQ KSKGPKKTAKSKKKKPLKK(K)TYTCAITTVKATETKAGKWSRWE

Caspase 5

(wt)ORF (SEQ ID NO: 30)

MFKGILQSGLDNFVINHMLKNNVAGQTSIQTLVPNTDQKSTSVKKDNHKKKTVKMLEYLGKDVLHGVFNYLAKHDVLTLKEEEKKKYY DAKIEDKALILVDSLRKNRVAHQMFTQTLLNMDQKITSVKPLLQIEAGPPESAESTNILKLCPREEFLRLCKKNHDEIYPIKKREDRRRLA LIICNTKFDHLPARNGAHYDIVGMKRLLQGLGYTVVDEKNLTARDMESVLRAFAARPEHKSSDSTFLVLMSHGILEGICGTAHKKKKPD VLLYDTIFQIFNNRNCLSLKDKPKVIIVQACRGEKHGELWVRDSPASLAVISSQSSENLEADSVCKIHEEKDFIAFCSSTPHNVSWRDR TRGSIFITELITCFQKYSCCCHLMEIFRKVQKSFEVPQAKAQMPTIERATLTRDFYLFPGN

(-1)ORF (SEQ ID NO: 31)

MFKGILQSGLDNFVINHMLKNNVAGQTSIQTLVPNTDQKSTSVKKDNHKKKQLRCWNTWAKMFFMVFLIIWQNTMF

(+1)/(-2)ORF (SEQ ID NO: 32)

MFKGILQSGLDNFVINHMLKNNVAGQTSIQTLVPNTDQKSTSVKKDNHKK(K)NS

AIM2

(wt)ORF (SEQ ID NO: 33)

MESKYKEILLLTGLDNITDEELDRFKFFLSDEFNIATGKLHTANRIQVATLMIQNAGAVSAVMKTIRIFQKLNYMLLAKRLQEEKEKVDKQ YKSVTKPKPLSQAEMSPAASAAIRNDVAKQRAAPKVSPHVKPEQKQMVAQQESIREGFQKRCLPVMVLKAKKPFTFETQEGKQEMF HATVATEKEFFFVKVFNTLLKDKFIPKRIIIIARYYRHSGFLEVNSASRVLDAESDQKVNVPLNIIRKAGETPKINTLQTQPLGTIVNGLFV VQKVTEKKKNILFDLSDNTGKMEVLGVRNEDTMKCKEGDKVRLTFFTLSKNGEKLQLTSGVHSTIKVIKAKKKT

(-1)ORF (SEQ ID NO: 34)

MESKYKEILLLTGLDNITDEELDRFKFFLSDEFNIATGKLHTANRIQVATLMIQNAGAVSAVMKTIRIFQKLNYMLLAKRLQEEKEKVDKQ YKSVTKPKPLSQAEMSPAASAAIRNDVAKQRAAPKVSPHVKPEQKQMVAQQESIREGFQKRCLPVMVLKAKKPFTFETQEGKQEMF HATVATEKEFFFVKVFNTLLKDKFIPKRIIIIARYYRHSGFLEVNSASRVLDAESDQKVNVPLNIIRKAGETPKINTLQTQPLGTIVNGLFV VQKVTEKKKNILFDLSDNTGKMEVLGVRNEDTMKCKEGDKVRLTFFTLSKNGEKLQLTSGVHSTIKVIKAKKKHREVKRTNSSQLV

(+1)/(-2)ORF (SEQ ID NO: 35)

MESKYKEILLLTGLDNITDEELDRFKFFLSDEFNIATGKLHTANRIQVATLMIQNAGAVSAVMKTIRIFQKLNYMLLAKRLQEEKEKVDKQ YKSVTKPKPLSQAEMSPAASAAIRNDVAKQRAAPKVSPHVKPEQKQMVAQQESIREGFQKRCLPVMVLKAKKPFTFETQEGKQEMF HATVATEKEFFFVKVFNTLLKDKFIPKRIIIIARYYRHSGFLEVNSASRVLDAESDQKVNVPLNIIRKAGETPKINTLQTQPLGTIVNGLFV VQKVTEKKKNILFDLSDNTGKMEVLGVRNEDTMKCKEGDKVRLTFFTLSKNGEKLQLTSGVHSTIKVIKAKK(K)*NIEK*

SLC23A1

(wt)ORF (SEQ ID NO: 36)

MMGIGKNTTSKSMEAGSSTEGKYEDEAKHPAFFTLPVVINGGATSSGEQDNEDTELMAIYTTENGIAEKSSLAETLDSTGSLDPQRS DMIYTIEDVPPWYLCIFLGLQHYLTCFSGTIAVPFLLADAMCVGYDQWATSQLIGTIFFCVGITTLLQTTFGCRLPLFQTSAFAFLAPARA ILSLDKWKCNTTDVSVANGTAELLHTEHIWYPRIREIQGAIIMSSLIEVVIGLLGLPGALLKYIGPLTITPTVALIGLSGFQAAGERAGKHW GIAMLTIFLVLLFSQYARNVKFPLPIYKSKKGWTAYKLQLFKMFPIILAILVSWLLCFIFTVTDVFPPDSTKYGFYARTDARQGVLLVAPW FKVPYPFQWGLPTVSAAGVIGMLSAVVASIIESIGDYYACARLSCAPPPPIHAINRGIFVEGLSCVLDGIFGTGNGSTSSSPNIGVLGITK VGSRRVIQCGAALMLALGMIGKFSALFASLPDPVLGALFCTLFGMITAVGLSNLQFIDLNSSRNLFVLGFSIFFGLVLPSYLRQNPLVTGI TGIDQVLNVLLTTAMFVGGCVAFILDNTIPGTPEERGIRKWKKGVGKGNKSLDGMESYNLPFGMNIIKKYRCFSYLPISPTFVGYTWK GLRKSDNSRSSDEDSQATG

(-1)ORF (SEQ ID NO: 37)

MMGIGKNTTSKSMEAGSSTEGKYEDEAKHPAFFTLPVVINGGATSSGEQDNEDTELMAIYTTENGIAEKSSLAETLDSTGSLDPQRS DMIYTIEDVPPWYLCIFLGLQHYLTCFSGTIAVPFLLADAMCVGYDQWATSQLIGTIFFCVGITTLLQTTFGCRLPLFQTSAFAFLAPARA ILSLDKWKCNTTDVSVANGTAELLHTEHIWYPRIREIQGAIIMSSLIEVVIGLLGLPGALLKYIGPLTITPTVALIGLSGFQAAGERAGKHW GIAMLTIFLVLLFSQYARNVKFPLPIYKSKKGWTAYKLQLFKMFPIILAILVSWLLCFIFTVTDVFPPDSTKYGFYARTDARQGVLLVAPW FKVPYPFQWGLPTVSAAGVIGMLSAVVASIIESIGDYYACARLSCAPPPPSTQ

(+1)/(-2)ORF (SEQ ID NO: 38)

MMGIGKNTTSKSMEAGSSTEGKYEDEAKHPAFFTLPVVINGGATSSGEQDNEDTELMAIYTTENGIAEKSSLAETLDSTGSLDPQRS DMIYTIEDVPPWYLCIFLGLQHYLTCFSGTIAVPFLLADAMCVGYDQWATSQLIGTIFFCVGITTLLQTTFGCRLPLFQTSAFAFLAPARA ILSLDKWKCNTTDVSVANGTAELLHTEHIWYPRIREIQGAIIMSSLIEVVIGLLGLPGALLKYIGPLTITPTVALIGLSGFQAAGERAGKHW GIAMLTIFLVLLFSQYARNVKFPLPIYKSKKGWTAYKLQLFKMFPIILAILVSWLLCFIFTVTDVFPPDSTKYGFYARTDARQGVLLVAPW FKVPYPFQWGLPTVSAAGVIGMLSAVVASIIESIGDYYACARLSCAPPP (P)HPRNKQGNFRGRPLLCS

ABCF1

(wt)ORF (SEQ ID NO: 39)

MPKAPKQQPPEPEWIGDGESTSPSDKVVKKGKKDKKIKKTFFEELAVEDKQAGEEEKVLKEKEQQQQQQQQQQKKKRDTRKGRR KKDVDDDGEEKELMERLKKLSVPTSDEEDEVPAPKPRGGKKTKGGNVFAALIQDQSEEEEEEEKHPPKPAKPEKNRINKAVSEEQQ PALKGKKGKEEKSKGKAKPQNKFAALDNEEEDKEEEIIKEKEPPKQGKEKAKKAEQMEYERQVASLKAANAAENDFSVSQAEMSSR QAMLENASDIKLEKFSISAHGKELFVNADLYIVAGRRYGLVGPNGKGKTTLLKHIANRALSIPPNIDVLLCEQEVVADETPAVQAVLRAD TKRLKLLEEERRLQGQLEQGDDTAAERLEKVYEELRATGAAAAEAKARRILAGLGFDPEMQNRPTQKFSGGWRMRVSLARALFMEP TLLMLDEPTNHLDLNAVIWLNNYLQGWRKTLLIVSHDQGFLDDVCTDIIHLDAQRLHYYRGNYMTFKKMYQQKQKELLKQYEKQEKKL KELKAGGKSTKQAEKQTKEALTRKQQKCRRKNQDEESQEAPELLKRPKEYTVRFTFPDPPPLSPPVLGLHGVTFGYQGQKPLFKNL DFGIDMDSRICIVGPNGVGKSTLLLLTGKLTPTHGEMRKNHRLKIGFFNQQYAEQLRMEETPTEYLQRGFNLPYQDARKCLGRFGLE SHAHTIQICKLSGGQKARVVFAELACREPDVLILDEPTNNLDIESIDALGEAINEYKGAVIVVSHDARLITETNCQLWVVEEQSVSQIDG DFEDYKREVLEALGEVMVSRPRE

(-1)ORF (SEQ ID NO: 40)

MPKAPKQQPPEPEWIGDGESTSPSDKVVKKGKKDKKIKKTFFEELAVEDKQAGEEEKVLKEKEQQQQQQQQQQKK<u>SEIPEKAGGR</u> RMWMMMEKRKSSWSVLRSSQCQPVMRRMKYPPQNPAEGRKPRVVMFLQP

(+1)/(-2)ORF (SEQ ID NO: 41)

MPKAPKQQPPEPEWIGDGESTSPSDKVVKKGKKDKKIKKTFFEELAVEDKQAGEEEKVLKEKEQQQQQQQQQKK(K)<u>ARYPKRQA</u> <u>EEGCG</u>

HSPC259

(wt)ORF (SEQ ID NO: 42)

SPDYFPQISSQFGTVEK

222

MEKIFISSSTKAEGKGISPFEAPINTQAPPEKGKEAVVQEPERSWFQTKEERKKEKIAKALQEFDLALRGKKKRKKFMKDAKKKGEMT AEERSQFEILKAQMFAERLAKRNRRAKRARAMPEEEPVRGPAKKQKQGKKSVFDEELTNTSKKALKQYRAGPSFEERKQLGLPHQR RGGNFKSNPDTRGGSSCRGLKKFMGAALKSLPCGKSSWLVCLFSICLKKKQKQKTTLWCGGMVRSYFPKHVCQSPFLLISFHMTIL NGSIFGKRE

(-1)ORF (SEQ ID NO: 43)

MÉKIFISSSTKAEGKGISPFEAPINTQAPPEKGKEAVVQEPERSWFQTKEERKKEKIAKALQEFDLALRGKKKRKKFMKDAKKKGEMT AEERSQFEILKAQMFAERLAKRNRRAKRARAMPEEPVRGPAKKQKQGKKSVFDEELTNTSKKALKQYRAGPSFEERKQLGLPHQR RGGNFKSNPDTRGGSSCRGLKKFMGAALKSLPCGKSSWLVCLFSICLKK*NKNKKQHFGVVVWYVAIFLSMSVNLPSC*

(+1)/(-2)ORF (SEQ ID NO: 44)

MEKIFISSSTKAEGKGISPFEAPINTQAPPEKGKEAVVQEPERSWFQTKEERKKEKIAKALQEFDLALRGKKKRKKFMKDAKKKGEMT AEERSQFEILKAQMFAERLAKRNRRAKRARAMPEEEPVRGPAKKQKQGKKSVFDEELTNTSKKALKQYRAGPSFEERKQLGLPHQR RGGNFKSNPDTRGGSSCRGLKKFMGAALKSLPCGKSSWLVCLFSICLKK(K)<u>TKTKNNTLVWWYGT</u>

Bax

(wt)ORF (SEQ ID NO: 45)

MDGSGEQPRGGGPTSSEQIMKTGALLLQGFIQDRAGRMGGEAPELALDPVPQDASTKKLSECLKRIGDELDSNMELQRMIAAVDTD SPREVFRVAADMFSDGNFNWGRVVALFYFASKLVLKALCTKVPELIRTIMGWTLDFLRERLLGWIQDQGGWDGLLSYFGTPTWQT VTIFVAGVLTASLTIWKKMG

(-1)ORF (SEQ ID NO: 46)

MDGSGEQPRGGGPTSSEQIMKTGALLLQGFIQDRAGRMGGRHPSWPWTRCLRMRPPRS

(+1)/(-2)ORF (SEQ ID NO: 47)

MDGSGEQPRGGGPTSSEQIMKTGALLLQGFIQDRAGRMGG(G)GTRAGPGPGASGCVHQEAERVSQAHRGRTGQ

TCF6L1

(wt)ORF (SEQ ID NO: 48)

MAFLRSMWGVLSALGRSGAELCTGCGSRLRSPFSFVYLPRWFSSVLASCPKKPVSSYLRFSKEQLPIFKAQNPDAKTTELIRRIAQR WRELPDSKKKIYQDAYRAEWQVYKEEISRFKEQLTPSQIMSLEKEIMDKHLKRKAMTKKKELTLLGKPKRPRSAYNVYVAERFQEAK GDSPQEKLKTVKENWKNLSDSEKELYIQHAKEDETRYHNEMKSWEEQMIEVGRKDLLRRTIKKQRKYGAEEC

(-1)ORF (SEQ ID NO: 49)

MÁFLRSMWGVLSALGRSGAELCTGCGSRLRSPFSFVYLPRWFSSVLASCPKKPVSSYLRFSKEQLPIFKAQNPDAKTTELIRRIAQR WRELPDSKKKIYQDAYRAEWQVYKEEISRFKEQLTPSQIMSLEKEIMDKHLKRKAMTKKKS

(+1)/(-2)ORF (SEQ ID NO: 50)

MAFLRSMWGVLSALGRSGAELCTGCGSRLRSPFSFVYLPRWFSSVLASCPKKPVSSYLRFSKEQLPIFKAQNPDAKTTELIRRIAQR WRELPDSKKKIYQDAYRAEWQVYKEEISRFKEQLTPSQIMSLEKEIMDKHLKRKAMTKK(K)*RVNTAWKTKKTSFSL*

FTL3I

(wt)ORF (SEQ ID NO: 51)

MTVLAPÄWSPTTYLLLLĹLLSSGLSGTQDCSFQHSPISSDFAVKIRELSDYLLQDYPVTVASNLQDEELCGGLWRLVLAQRWMERLKT VAGSKMQGLLERVNTEIHFVTKCAFQPPPSCLRFVQTNISRLLQETSEQLVALKPWITRQNFSRCLELQCQPDSSTLPPPWSPRPLE ATAPTAPQPPLLLLLLPVGLLLLAAAWCLHWQRTRRRTPRPGEQVPPVPSPQDLLLVEH

(-1)ORF (SEQ ID NO: 52)

MTVLAPAWSPTTYLLLLLLSSGLSGTQDCSFQHSPISSDFAVKIRELSDYLLQDYPVTVASNLQDEELCGGLWRLVLAQRWMERLKT VAGSKMQGLLERVNTEIHFVTKCAFQ

PPPAVFASSRPTSPASCRRPPSSWWR

(+1)/(-2)ORF (SEQ ID NO: 53)

MTVLAPAWSPTTYLLLLLLLSSGLSGTQDCSFQHSPISSDFAVKIRELSDYLLQDYPVTVASNLQDEELCGGLWRLVLAQRWMERLKT VAGSKMQGLLERVNTEIHFVTKCAFQ

PP(P)QLSSLRPDQHLPPPAGDLRAAGGAEALDHSPELLPVPGAAVSAR

OGT

(wt)ORF (SEQ ID NO: 54)

MLQGHFWLVREGIMISPSSPPPPNLFFFPLQIFPFFTSFPSHLLSLTPPKACYLKAIETQPNFAVAWSNLGCVFNAQGEIWLAIHHFE KAVTLDPNFLDAYINLGNVLKEARIFDRAVAAYLRALSLSPNHAVVHGNLACVYYEQGLIDLAIDTYRRAIELQPHFPDAYCNLANALKE KGSVAEAEDCYNTALRLCPTHADSLNNLANIKREQGNIEEAVRLYRKALEVFPEFAAAHSNLASVLQQQGKLQEALMHYKEAIRISPTF ADAYSNMGNTLKEMQDVQGALQCYTRAIQINPAFADAHSNLASIHKDSGNIPEAIASYRTALKLKPDFPDAYCNLAHCLQIVCDWTDY DERMKKLVSIVADQLEKNRLPSVHPHHSMLYPLSHGFRKAIAERHGNLCLDKINVLHKPPYEHPKDLKLSDGRLRVGYVSSDFGNHPT SHLMQSIPGMHNPDKFEVFCYALSPDDGTNFRVKVMAEANHFIDLSQIPCNGKAADRIHQDGIHILVNMNGYTKGARNELFALRPAPI QAMWLGYPGTSGALFMDYIITDQETSPAEVAEQYSEKLAYMPHTFFIGDHANMFPHLKKKAVIDFKSNGHIYDNRIVLNGIDLKAFLDS LPDVKIVKMKCPDGGDNADSSNTALNMPVIPMNTIAEAVIEMINRGQIQITINGFSISNGLATTQINNKAATGEEVPRTIIVTTRSQYGLP EDAIVYCNFNQLYKIDPSTLQMWANILKRVPNSVLWLLRFPAVGEPNIQQYAQNMGLPQNRIIFSPVAPKEEHVRRGQLADVCLDTPL CNGHTTGMDVLWAGTPMVTMPGETLASRVAASQLTCLGCLELIAKNRQEYEDIAVKLGTDLEYLKKVRGKVWKQRISSPLFNTKQYT MELERLYLQMWEHYAAGNKPDHMIKPVEVTESA

(-1)ORF (SEQ ID NO: 55)

MLQGHFWLVREGIMISPSSPPPPNLFFSLYKFSPFPLPPFPPIFFH

(+1)/(-2)ORF (SEQ ID NO: 56)

MLQGHFWLVREGIMISPSSPPPPNLFF(F)PFTNFPLSLYLLSLPSSFINPS

ELAVL3

(wt)ORF (SEQ ID NO: 57)

MESQVGGGPAGRPAQRPLLGTNGATDDSKTNLIVNYLPQNMTQDEFKSLFGSIGDIESCKLVRDKITGQSLGYGFVNYSDPNDADKA INTLNGLKLQTKTIKVSYARPSSASIRDANLYVSGLPKTMSQKEMEQLFSQYGRIITSRILVDQVTGVSRGVGFIRFDKRIEAEEAIKGLN GQKPLGAREPITVKFANNPSQKTGQALLTHLYQSSARRYAGPLHHQTQRFRLDNLLNMAYAVKRFSPIAIDGMSGLAGVGLSGGAAG GWCIFVYNLSPEPDQSVLWQLFGPFGAVTNVKVIRDFTTNKCKGFGFMTMTNYDEAAMAIASLNGYRLGQRVLQVSFKTSKQHKA

(-1)ORF (SEQ ID NO: 58)

MESQVGGARPAGLPNGHSLVQMEPLTTARPTSSSTTCPRT

(+1)/(-2)ORF (SEQ ID NO: 59)

MESQVGG(G)PGRPACPTATPWYKWSH

MAC30X

(wt)ORF (SEQ ID NO: 60)

LFSHQRVQAQPTDYGGSFTRRCVEWLLGLYFLSHIPITLFMDLQAVVPRELYPVEFRNLLKWYAKEFKDPLLQEPPAWFKSFLFCELV FQLPFFPIATYAFLKGSCKWIRTPAIIYSVHTMTTLILILSTFLFEDFSKASGFKGQRPETLHERLTLVSVYAPYLLIPFILLIFMLRSPYYKY EEKRKKK

(-1)ORF (SEQ ID NO: 61)

LFSHQRVQAQPTDYGGSFTRRCVEWLLGLYFLSHIPITLFMDLQAVVPRELYPVEFRNLLKWYAKEFKDPLLQEPPAWFKSFLFCELV FQLPFFPIATYAFLKGSCKWIRTPAIIYSVHTMTTLILLSTFLFEDFSKASGFKGQRPETLHERLTLVSVYAPYLLIPFILLIFMLRSPYYKY EEKRKKNEGNNHWPRVEMPTGWLLVGYIQEHCSEPTSSAAFETLAAMHKSKMVSGTMSNPHLLPFFFFF

(+1)/(-2)ORF (SEQ ID NO: 62)

LFSHQRVQAQPTDYGGSFTRRCVEWLLGLYFLSHIPITLFMDLQAVVPRELYPVEFRNLLKWYAKEFKDPLLQEPPAWFKSFLFCELV FQLPFFPIATYAFLKGSCKWIRTPAIIYSVHTMTTLILILSTFLFEDFSKASGFKGQRPETLHERLTLVSVYAPYLLIPFILLIFMLRSPYYKY EEKRKK(K)<u>MKETTTGPG</u>

SLC4A3

(wt)ORF (SEQ ID NO: 63)

MAŃGVIPPPGGASPLPQVRVPLEEPPLSPDVEEEDDDLGKTLAVSRFGDLISKPPAWDPEKPSRSYSERDFEFHRHTSHHTHHPLSA RLPPPHKLRRLPPTSARHTRKRKKEKTSAPPSEGTPPIQEEGGAGVDEEEEEEEEEGESEAEPVEPPPSGTPQKAKFSIGSDEDD SPGLPGRAAVTKPLPSVGPHTDKSPQHSSSSPSPRARASRLAGEKSRPWSPSASYDLRERLCPGSALGNPGGPEQQVPTDEAEAQ MLGSADLDDMKSHRLEDNPGVRRHLVKKPSRTQGGRGSPSGLAPILRRKKKKKLDRRPHEVFVELNELMLDRSQEPHWRETARW IKFEEDVEETERWGKPHVASLSFRSLLELRRTIAHGAALLDLEQTTLPGIAHLVVETMIVSDQIRPEDRASVLRTLLLKHSHPNDDKDS GFFPRNPSSSSMNSVLGNHHPTPSHGPDGAVPTMADDLGEPAPLWPHDPDAKEKPLHMPGGDGHRGKSLKLLEKIPEDAEATVVL VGCVPFLEQPAAAFVRLNEAVLLESVLEVPVPVRFLFVMLGPSHTSTDYHELGRSIATLMSDKLFHEAAYQADDRQDLLSAISEFLDG SIVIPPSEVEGRDLLRSVAAFQRELLRKRREREQTKVEMTTRGGYTAPGKELSLELGGSEATPEDDPLLRTGSVFGGLVRDVRRRYP HYPSDLRDALHSQCVAAVLFIYFAALSPAITFGGLLGEKTEGLMGVSELIVSTAVLGVLFSLLGAQPLLVVGFSGPLLVFEEAFFKFCRA QDLEYLTGRVWVGLWLVVFVLALVAAEGSFLVRYISPFTQEIFAFLISLIFIYETFYKLYKVFTEHPLLPFYPPEGALEGSLAAGLEPNGS ALPPTEGPPSPRNQPNTALLSLILMLGTFFIAFFLRKFRNSRFLGGKARRIIGDFGIPISILVMVLVDYSITDTYTQKLTVPTGLSVTSPDK RSWFIPPLGSARPFPPWMMVAAAVPALLVLILIFMETQITALIVSQKARRILKGSGFHLDLLLIGSLGGLCGLFGLPWLTAATVRSVTHV NALTVMRTAIAPGDKPQIQEVREQRVTGVLIASLVGLSIVMGAVLRRIPLAVLFGIFLYMGVTSLSGIQLSQRLLLILMPAKHHPEQPYVT KVKTWRMHLFTCIQLGCIALLWVVKSTAASLAFPFLLLLTVPLRHCLLPRLFQDRELQALDSEDAEPNFDEDGQDEYNELHMPV

(-1)ORF (SEQ ID NO: 64)

MANGVIPPPGGASPLPQVRVPLEEPPLSPDVEEEDDDLGKTLAVSRFGDLISKPPAWDPEKPSRSYSERDFEFHRHTSHHTHHPLSA RLPPPHKLRRLPPTSARHTRRKRKKEKTSAPPSEGTPPIQEEGGAGVDEEEEEEEEEEEEEEEAEPVEPPPQGPHRRQSSPLEVTRM TVQASLGGLLSPSPCPRWAHTLTRAPSTPAAPPAPGPGPPDSLGRKAGPGAHRPVMTCGSDCAQAVPWATQVVQSSRCPQMRRR PRCWVLQTWTT

(+1)/(-2)ORF (SEQ ID NO: 65)

MANGVIPPPGGASPLPQVRVPLEEPPLSPDVEEEDDDLGKTLAVSRFGDLISKPPAWDPEKPSRSYSERDFEFHRHTSHHTHHPLSA RLPPPHKLRRLPPTSARHTRRKRKKEKTSAPPSEGTPPIQEEGGAGVDEEEEEEEEEEEEEEPVEPP(P)LRDPTEGKVLHWK

PRKDC

(wt)ORF (SEQ ID NO: 66)

MAGSGAGVRCSLLRLQÉTLSAADRCGAALAGHQLIRGLGQECVLSSSPAVLALQTSLVFSRDFGLLVFVRKSLNSIEFRECREEILKFL CIFLEKMGQKIAPYSVEIKNTCTSVYTKDRAAKCKIPALDLLIKLLQTFRSSRLMDEFKIGELFSKFYGELALKKKIPDTVLEKVYELLGLL GEVHPSEMINNAENLFRAFLGELKTQMTSAVREPKLPVLAGCLKGLSSLLCNFTKSMEEDPQTSREIFNFVLKAIRPQIDLKRYAVPSA GLRLFALHASQFSTCLLDNYVSLFEVLLKWCAHTNVELKKAALSALESFLKQVSNMVAKNAEMHKNKLQYFMEQFYGIIRNVDSNNKE LSIAIRGYGLFAGPCKVINAKDVDFMYVELIQRCKQMFLTQTDTGDDRVYQMPSFLQSVASVLLYLDTVPEVYTPVLEHLVVMQIDSFP QYSPKMQLVCCRAIVKVFLALAAKGPVLRNCISTVVHQGLIRICSKPVVLPKGPESESDHRASGEVRTGKWKVPTYKDYVDLFRHLL SSDQMMDSILADEAFFSVNSSSESLNHLLYDEFVKSVLKIVEKLDLTLEIQTVGEQENGDEAPGVWMIPTSDPAANLHPAKPKDFSAFI NLVEFCREILPEKQAEFFEPWVYSFSYELILQSTRLPLISGFYKLLSITVRNAKKIKYFEGS

(-1)ORF (SEQ ID NO: 67)

MÁGSGÁGVRCSLLRLQÉTLSAADRCGAALAGHQLIRGLGQECVLSSSPAVLALQTSLVFSRDFGLLVFVRKSLNSIEFRECREEILKFL CIFLEKMGQKIAPYSVEIKNTCTSVYTKDRAAKCKIPALDLLIKLLQTFRSSRLMDEFKIGELFSKFYGELALKKK<u>YQIQF</u>

(+1)/(-2)ORF (SEQ ID NO: 68

MAGSGAGVRCSLLRLQETLSAADRCGAALAGHQLIRGLGQECVLSSSPAVLALQTSLVFSRDFGLLVFVRKSLNSIEFRECREEILKFL CIFLEKMGQKIAPYSVEIKNTCTSVYTKDRAAKCKIPALDLLIKLLQTFRSSRLMDEFKIGELFSKFYGELALKK(K)<u>NTRYSFRKSI</u>

IIV/RAG

(wt)ORF) (SEQ ID NO: 69)

MSÁSASÝGGPVPQPPPGPAAALPPGSAARALHVELPSQQRRLRHLRNIAARNIVNRNGHQLLDTYFTLHLCSTEKIYKEFYRSEVIKN SLNPTWRSLDFGIMPDRLDTSVSCFVVKIWGGKENIYQLLIEWKVCLDGLKYLGQQIHARNQNEIIFGLNDGYYGAPFEHKGYSNAQK TILLQVDQNCVRNSYDVFSLLRLHRAQCAIKQTQVTVQKIGKEIEEKLRLTSTSNELKKKSECLQLKILVLQNELERQKKALGREVALLH KQQIALQDKGSAFSAEHLKLQLQKESLNELRKECTAKRELFLKTNAQLTIRCRQLLSELSYIYPIDLNEHKDYFVCGVKLPNSEDFQAK DDGSIAVALGYTAHLVSMISFFLQVPLRYPIIHKGSRSTIKDNINDKLTEKEREFPLYPKGGEKLQFDYGVYLLNKNIAQLRYQHGLGTP DLRQTLPNLKNFMEHGLMVRCDRHHTSSAIPVPKRQSSIFGGADVGFSGGIPSPDKGHRKRASSENERLQYKTPPPSYNSALAQPVT TVPSMGETERKITSLSSSLDTSLDFSKENKKKGEDLVGSLNGGHANVHPSQEQGEALSGHRATVNGTLLPSEQAGSASVQLPGEFH PVSEAELCCTVEQAEEIIGLEAQVSPQVIS

(-1)ORF (SEQ ID NO: 70)

MSASASVGGPVPQPPPGPAAALPPGSAARALHVELPSQQRRLRHLRNIAARNIVNRNGHQLLDTYFTLHLCSTEKIYKEFYRSEVIKN SLNPTWRSLDFGIMPDRLDTSVSCFVVKIWGGKENIYQLLIEWKVCLDGLKYLGQQIHARNQNEIIFGLNDGYYGAPFEHKGYSNAQK TILLQVDQNCVRNSYDVFSLLRLHRAQCAIKQTQVTVQKIGKEIEEKLRLTSTSNELKKK<u>VVACS</u>

(+1)/(-2)ORF (SEQ ID NO: 71)

MSASÁSVGGPVPQPPPGPÁAALPPGSAARALHVELPSQQRRLRHLRNIAARNIVNRNGHQLLDTYFTLHLCSTEKIYKEFYRSEVIKN SLNPTWRSLDFGIMPDRLDTSVSCFVVKIWGGKENIYQLLIEWKVCLDGLKYLGQQIHARNQNEIIFGLNDGYYGAPFEHKGYSNAQK TILLQVDQNCVRNSYDVFSLLRLHRAQCAIKQTQVTVQKIGKEIEEKLRLTSTSNELKKK<u>(K)</u>

MSH3

(wt)ORF (SEQ ID NO: 72)

MSRRKPASGGLAASSSÁPARQAVLSRFFQSTGSLKSTSSSTGAADQVDPGAAAAAAAAAAAAAPPAPPAPAFPPQLPPHVATEIDRR KKRPLENDGPVKKKVKVQQKEGGSDLGMSGNSEPKKCLRTRNVSKSLEKLKEFCCDSALPQSRVQTESLQERFAVLPKCTDFDDI SLLHAKNAVSSEDSKRQINQKDTTLFDLSQFGSSNTSHENLQKTASKSANKRSKSIYTPLELQYIEMKQQHKDAVLCVECGYKYRFFG EDAEIAARELNIYCHLDHNFMTASIPTHRLFVHVRRLVAKGYKVGVVKQTETAALKAIGDNRSSLFSRKLTALYTKSTLIGEDVNPLIKLD DAVNVDEIMTDTSTSYLLCISENKENVRDKKKGNIFIGIVGVQPATGEVVFDSFQDSASRSELETRMSSLQPVELLLPSALSEQTEALIH RATSVSVQDDRIRVERMDNIYFEYSHAFQAVTEFYAKDTVDIKGSQIISGIVNLEKPVICSLAAIIKYLKEFNLEKMLSKPENFKQLSSKM EFMTINGTTLRNLEILQNQTDMKTKGSLLWVLDHTKTSFGRRKLKKWVTQPLLKLREINARLDAVSEVLHSESSVFGQIENHLRKLPDI GRGLCSIYHKKCSTQEFFLIVKTLYHLKSEFQAIIPAVNSHIQSDLLRTVILEIPELLSPVEHYLKILNEQAAKVGDKTELFKDLSDFPLIKK RKDEIQGVIDEIRMHLQEIRKILKNPSAQYVTVSGQEFMIEIKNSAVSCIPTDWVKVGSTKAVSRFHSPFIVENYRHLNQLREQLVLDCS AEWLDFLEKFSEHYHSLCKAVHHLATVDCIFSLAKVAKQGDYCRPTVQEERKIVIKNGRHPVIDVLLGEQDQYVPNNTDLSEDSERVM ITGPNMGGKSSYIKQVALITIMAQIGSYVPAEEATIGIVDGIFTRMGAADNIYKGRSTFMEELTDTAEIIRKATSQSLVILDELGRGTSTH DGIAIAYATLEYFIRDVKSLTLFVTHYPPVCELEKNYSHQVGNYHMGFLVSEDESKLDPGTAEQVPDFVTFLYQITRGIAARSYGLNVA KLADVPGEILKKAAHKSKELEGLINTKRKRLKYFAKLWTMHNAQDLQKWTEEFNMEETQTSLLH

(-1)ORF (SEQ ID NO: 73)

MŚRRKPASGGLAASSSÁPARQAVLSRFFQSTGSLKSTSSSTGAADQVDPGAAAAAAAAAAAAPPAPPAPAFPPQLPPHVATEIDRR KKRPLENDGPVKKKVKKVQQKEGGSDLGMSGNSEPKKCLRTRNVSKSLEKLKEFCCDSALPQSRVQTESLQERFAVLPKCTDFDDI SLLHAKNAVSSEDSKRQINQKDTTLFDLSQFGSSNTSHENLQKTASKSANKRSKSIYTPLELQYIEMKQQHKDAVLCVECGYKYRFFG EDAEIAARELNIYCHLDHNFMTASIPTHRLFVHVRRLVAKGYKVGVVKQTETAALKAIGDNRSSLFSRKLTALYTKSTLIGEDVNPLIKLD DAVNVDEIMTDTSTSYLLCISENKENVRDKK*RATFLLALWECSLPQARLCLIVSRTLLLVQS*

(+1)/(-2)ORF (SEQ ID NO: 74)

MSRRKPASGGLAASSSAPARQAVLSRFFQSTGSLKSTSSSTGAADQVDPGAAAAAAAAAAAAAPPAPPAPAFPPQLPPHVATEIDRR KKRPLENDGPVKKKVKVQQKEGGSDLGMSGNSEPKKCLRTRNVSKSLEKLKEFCCDSALPQSRVQTESLQERFAVLPKCTDFDDI SLLHAKNAVSSEDSKRQINQKDTTLFDLSQFGSSNTSHENLQKTASKSANKRSKSIYTPLELQYIEMKQQHKDAVLCVECGYKYRFFG EDAEIAARELNIYCHLDHNFMTASIPTHRLFVHVRRLVAKGYKVGVVKQTETAALKAIGDNRSSLFSRKLTALYTKSTLIGEDVNPLIKLD DAVNVDEIMTDTSTSYLLCISENKENVRDKK(K)GQHFYWHCGSAACHRRGCV

ACVR2.

(wt) ORF (SEQ ID NO: 107)

MGAAAKLAFA VFLISCSSGA ILGRSETQEC LFFNANWEKD RTNQTGVEPC YGDKDKRRHC FATWKNISGS IEIVKQGCWL DDINCYDRTD CVEKKDSPEV YFCCCEGNMC NEKFSYFPEM EVTQPTSNPV TPKPPYYNIL LYSLVPLMLI AGIVICAFWV

YRHHKMAYPP VLVPTQDPGP PPPSPLLGLK PLQLLEVKAR GRFGCVWKAQ LLNEYVAVKI FPIQDKQSWQ NEYEVYSLPG MKHENILQFI GAEKRGTSVD VDLWLITAFH EKGSLSDFLK ANVVSWNELC HIAETMARGL AYLHEDIPGL KDGHKPAISH RDIKSKNVLL KNNLTACIAD FGLALKFEAG KSAGDTHGQV GTRRYMAPEV LEGAINFQRD AFLRIDMYAM GLVLWELASR CTAADGPVDE YMLPFEEEIG QHPSLEDMQE VVVHKKKRPV LRDYWQKHAG MAMLCETIEE CWDHDAEARL SAGCVGERIT QMQRLTNIIT TEDIVTVVTM VTNVDFPPKE SSL*

A8, Pos. 451: -1 ORF (Mut.rate 16.3%) (SEQ ID NO: 108)

MGAAAKLAFA VFLISCSSGA ILGRSETQEC LFFNANWEKD RTNQTGVEPC YGDKDKRRHC FATWKNISGS IEIVKQGCWL DDINCYDRTD CVEK*KTALKY IFVAVRAICV MKSFLIFRRW KSHSPLQIQL HLSHPITTSC SIPWCHLC***

A8 Pos. 1476: -1 (Mut.rate 81.6%) (SEQ ID NO: 109)

MGAAKLAFA VFLISCSSGA ILGRSETQEC LFFNANWEKD RTNQTGVEPC YGDKDKRRHC FATWKNISGS IEIVKQGCWL DDINCYDRTD CVEKKDSPEV YFCCCEGNMC NEKFSYFPEM EVTQPTSNPV TPKPPYYNIL LYSLVPLMLI AGIVICAFWV YRHHKMAYPP VLVPTQDPGP PPPSPLLGLK PLQLLEVKAR GRFGCVWKAQ LLNEYVAVKI FPIQDKQSWQ NEYEVYSLPG MKHENILQFI GAEKRGTSVD VDLWLITAFH EKGSLSDFLK ANVVSWNELC HIAETMARGL AYLHEDIPGL KDGHKPAISH RDIKSKNVLL KNNLTACIAD FGLALKFEAG KSAGDTHGQV GTRRYMAPEV LEGAINFQRD AFLRIDMYAM GLVLWELASR CTAADGPVDE YMLPFEEEIG QHPSLEDMQE VVVHKK*RGLF**

FLJ11053, A11 Pos. 1695, Mut.rate 52.2%

wt ORF (SEQ ID NO: 110)

MVLRKLSKKD VTTKLKAMQE FGTMCTERDT ETVKGVLPYW PRIFCKISLD HDRRVREATQ QAFEKLTLKV KKQLAPYLKS LMGYWLMAQC DTYTPAAFAA KDAFEAAFPP SKQPEAIAFC KDEITSVLQD HLIKETPDTL SDPQTVPEEE REAKFYRVVT CSLLALKRLL CLLPDNELDS LEEKFKSLLS QNKFWKYGKH SVPQIRSAYF ELVSALCQRI PQLMKEEASK VSPSVLLSID DSDPIVCPAL WEAVLYTLTI IEDCWLHVNA KKSVFPKLST VIREGGRGLA TVIYPYLLPF ISKLPHSITN PKLDFFKNFL TSLVAGLSTE RTKTSSSESS AVISAFYECL RFIMQQNLGE EEIEQMLVND QLIPFIDAVL KDPGLQHGQL FNHLAETLSS WEAKADTEKD EKTAHNLENV LIHFWERLSE ICVAKISEPE ADVESVLGVS NLLQVLQKPK SSLKSSKKKN GKVRFADEIL ESNKENEKCV SSEGEKIEGW ELTTEPSLTH NSSGLLSPLR KKPLEDLVCK LADISINYVN ERKSEQHLRF LSTLLDSFSS SRVFKMLLGD EKQSIVQAKP LEIAKLVQKN PAVQFLYQKL IGWLNEDQRK DFGFLVDILY SALRCCDNDM

-1 ORF (SEQ ID NO: 111)

MVLRKLSKKD VTTKLKAMQE FGTMCTERDT ETVKGVLPYW PRIFCKISLD HDRRVREATQ QAFEKLTLKV KKQLAPYLKS LMGYWLMAQC DTYTPAAFAA KDAFEAAFPP SKQPEAIAFC KDEITSVLQD HLIKETPDTL SDPQTVPEEE REAKFYRVVT CSLLALKRLL CLLPDNELDS LEEKFKSLLS QNKFWKYGKH SVPQIRSAYF ELVSALCQRI PQLMKEEASK VSPSVLLSID DSDPIVCPAL WEAVLYTLTT IEDCWLHVNA KKSVFPKLST VIREGGRGLA TVIYPYLLPF ISKLPHSITN PKLDFFKNFL TSLVAGLSTE RTKTSSSESS AVISAFYECL RFIMQQNLGE EEIEQMLVND QLIPFIDAVL KDPGLQHGQL FNHLAETLSS WEAKADTEKD EKTAHNLENV LIHFWERLSE ICVAKISEPE ADVESVLGVS NLLQVLQKPK SSLKSSKKKM VRLDLLMRYL KAIKRMKNVY LQKERRLKAG N*

-2 ORF (SEQ ID NO: 112)

MVLRKLSKKD VTTKLKAMQE FGTMCTERDT ETVKGVLPYW PRIFCKISLD HDRRVREATQ QAFEKLTLKV KKQLAPYLKS LMGYWLMAQC DTYTPAAFAA KDAFEAAFPP SKQPEAIAFC KDEITSVLQD HLIKETPDTL SDPQTVPEEE REAKFYRVVT CSLLALKRLL CLLPDNELDS LEEKFKSLLS QNKFWKYGKH SVPQIRSAYF ELVSALCQRI PQLMKEEASK VSPSVLLSID DSDPIVCPAL WEAVLYTLTT IEDCWLHVNA KKSVFPKLST VIREGGRGLA TVIYPYLLPF ISKLPHSITN PKLDFFKNFL TSLVAGLSTE RTKTSSSESS AVISAFYECL RFIMQQNLGE EEIEQMLVND QLIPFIDAVL KDPGLQHGQL FNHLAETLSS WEAKADTEKD EKTAHNLENV LIHFWERLSE ICVAKISEPE ADVESVLGVS NLLQVLQKPK SSLKSSKKK<u>W</u>*

+ 1 ORF (SEQ ID NO: 113)

MVLRKLSKKD VTTKLKAMQE FGTMCTERDT ETVKGVLPYW PRIFCKISLD HDRRVREATQ QAFEKLTLKV KKQLAPYLKS LMGYWLMAQC DTYTPAAFAA KDAFEAAFPP SKQPEAIAFC KDEITSVLQD HLIKETPDTL SDPQTVPEEE REAKFYRVVT CSLLALKRLL CLLPDNELDS LEEKFKSLLS QNKFWKYGKH SVPQIRSAYF ELVSALCQRI PQLMKEEASK VSPSVLLSID DSDPIVCPAL WEAVLYTLTT IEDCWLHVNA KKSVFPKLST VIREGGRGLA TVIYPYLLPF ISKLPHSITN PKLDFFKNFL TSLVAGLSTE RTKTSSSESS AVISAFYECL RFIMQQNLGE EEIEQMLVND QLIPFIDAVL KDPGLQHGQL FNHLAETLSS WEAKADTEKD EKTAHNLENV LIHFWERLSE ICVAKISEPE ADVESVLGVS NLLQVLQKPK SSLKSSKKKK \underline{W}^*

KIAA1052, A11 Pos. 689, Mut.rate 42.2%

Wt ORF (SEQ ID NO: 114)

MAGRPLRIGD QLVLEEDYDE TYIPSEQEIL EFAREIGIDP IKEPELMWLA REGIVAPLPG EWKPCQDITG DIYYFNFANG QSMWDHPCDE HYRSLVIQER AKLSTSGAIK KKKKKKEKKD KKDRDPPKSS LALGSSLAPV HVPLGGLAPL RGLVDTPPSA LRGSQSVSLG SSVESGRQLG ELMLPSQGLK TSAYTKGLLG SIYEDKTALS LLGLGEETNE EDEEESDNQS VHSSSEPLRN LHLDIGALGG DFEYEESLRT SQPEEKKDVS LDSDAAGPPT PCKPSSPGAD SSLSSAVGKG RQGSGARPGL PEKEENEKSE PKICRNLVTP KADPTGSEPA KASEKEAPED TVDAGEEGSR REEAAKEPKK KASALEEGSS DASQELEISE HMKEPQLSDS IASDPKSFHG LDFGFRSRIS EHLDVDVLS PVLGGACRQA QQPLGIEDKD DSQSSQDELQ SKQSKGLEER YHRLSPPLPH EERAQSPPRS LATEEEPPQG PEGQPEWKEA EELGEDSAAS LSLQLSLQRE QAPSPPAACE KGKEQHSQAE ELGPGQEEAE DPEEKVAVSP TPPVSPEVRS TEPVAPPEQL SEAALKAMEE AVAQVLEQDQ RHLLESKQEK MQQLREKLCQ EEEEIIRLH QQKEQSLSSL RERLQKAIEE EEARMREEES QRLSWLRAQV QSSTQADEDQ IRAEQEASLQ KLREELESQQ KAERASLEQK

NRQMLEQLKE EIEASEKSEQ AALNAAKEKA LQQLREQLEG ERKEAVATLE KEHSAELERL CSSLEAKHRE VVSSLQKKIQ EAQQKEEAQL QKCLGQVEHR VHQKSYHVAG YEHELSSLLR EKRQEVEGEH ERRLDKMKEE HQQVMAKARE QYEAEERKQR AELLGHLTGE LERLQRAHER ELETVRQEQH KRLEDLRRRH REQERKLQDL ELDLETRAKD VKARLALLEV QEETARREKQ QLLDVQRQVA LKSEEATATH QQLEEAQKEH THLLQSNQQL REILDELQAR KLKLESQVDL LQAQSQQLQK HFSSLEAEAQ KKQHLLREVT VEENNASPHF EPDLHIEDLR KSLGTNQTKE VSSSLSQSKE DLYLDSLSSH NVWHLLSAEG VALRSAKEFL VQQTRSMRRR QTALKAAQQH WRHELASAQE VAKDPPGIKA LEDMRKNLEK ETRHLDEMKS AMRKGHNLLK KKEEKLNQLE SSLWEEASDE GTLGGSPTKK AVTFDLSDMD SLSSESSESF SPPHLDSTPS LTSRKIHGLS HSLRQISSQL SSVLSILDSL NPQSPPPLLA SMPAQLPPRD PKSTPTPTYY GSLARFSALS SATPTSTQWA WDSGQGPRLP SSVAQTVDDF LLEKWRKYFP SGIPLLSNSP TPLESRLGYM SASEQLRLLQ HSHSQVPEAG STTFQGIIEA NRRWLERVKN DPRLPLFSST PKPKATLSLL QLGLDEHNRV KVYRF*

- 1 ORF (SEQ ID NO: 115)

MAGRPLRIGD QLVLEEDYDE TYIPSEQEIL EFAREIGIDP IKEPELMWLA REGIVAPLPG EWKPCQDITG DIYYFNFANG QSMWDHPCDE HYRSLVIQER AKLSTSGAIK KKKKK<u>RKRKT RRTETPPKVR WPWVPH</u>**

- 20RF (SEQ ID NO: 116)

MAGRPLRIGD QLVLEEDYDE TYIPSEQEIL EFAREIGIDP IKEPELMWLA REGIVAPLPG EWKPCQDITG DIYYFNFANG QSMWDHPCDE HYRSLVIQER AKLSTSGAIK KKKKK<u>GKERQ EGQRPPQKFA GLGFLISPSS CSSWGPGSFT RSCGYPTLCS SWISKREPGE LSGVWTSAWR THAAFTGSQD LCLYKGSLGLHI*</u>

+ 1 ORF (SEQ ID NO: 117)

MAGRPLRIGD QLVLEEDYDE TYIPSEQEIL EFAREIGIDP IKEPELMWLA REGIVAPLPG EWKPCQDITG DIYYFNFANG QSMWDHPCDE HYRSLVIQER AKLSTSGAIK KKKKKK<u>GKER QEGQRPPQKF AGLGFLISPS SCSSWGPGSF TRSCGYPTLC SSWISKREPG ELSGVWTSAW RTHAAFTGSQ DLCLYKGSLG LHI*</u>